RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/550,280
Source: PCT
Date Processed by STIC: 10/06/2005

ENTERED

PCT

RAW SEQUENCE LISTING DATE: 10/06/2005
PATENT APPLICATION: **US/10/550,280** TIME: 11:41:37

Input Set : **A:\Sequence Listing 2005_1500A.txt**
Output Set: **N:\CRF4\10062005\J550280.raw**

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      3 <110> APPLICANT: NIPPON MEDICAL SCHOOL FOUNDATION
      5 <120> TITLE OF INVENTION: Cell death-inducing fusion gene specifically acting on
cancer and
      6       gene product thereof
      8 <130> FILE REFERENCE: PH-2011-PCT
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/550,280
C--> 11 <141> CURRENT FILING DATE: 2005-09-23
     13 <150> PRIOR APPLICATION NUMBER: JP2003/081337
     14 <151> PRIOR FILING DATE: 2003-03-24
     16 <160> NUMBER OF SEQ ID NOS: 27
     17 <170> SOFTWARE: PatentIn Ver. 2.1
     19 <210> SEQ ID NO: 1
     20 <211> LENGTH: 579
     21 <212> TYPE: DNA
     22 <213> ORGANISM: Homo sapiens
     24 <220> FEATURE:
     25 <221> NAME/KEY: CDS
     26 <222> LOCATION: (1)..(579)
     28 <400> SEQUENCE: 1
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     30 Met Asp Gly Ser Gly Glu Gln Pro Arg Gly Gly Gly Pro Thr Ser Ser
     31   1               5                  10                  15
     33 gag cag atc atg aag aca ggg gcc ctt ttg ctt cag ggt ttc atc cag   96
     34 Glu Gln Ile Met Lys Thr Gly Ala Leu Leu Leu Gln Gly Phe Ile Gln
     35              20                  25                  30
     37 gat cga gca ggg cga atg ggg ggg gag gca ccc gag ctg gcc ctg gac   144
     38 Asp Arg Ala Gly Arg Met Gly Gly Glu Ala Pro Glu Leu Ala Leu Asp
     39          35                  40                  45
     41 ccg gtg cct cag gat gcg tcc acc aag aag ctg agc gag tgt ctc aag   192
     42 Pro Val Pro Gln Asp Ala Ser Thr Lys Lys Leu Ser Glu Cys Leu Lys
     43      50                  55                  60
     45 cgc atc ggg gac gaa ctg gac agt aac atg gag ctg cag agg atg att   240
     46 Arg Ile Gly Asp Glu Leu Asp Ser Asn Met Glu Leu Gln Arg Met Ile
     47  65                  70                  75                  80
     49 gcc gcc gtg gac aca gac tcc ccc cga gag gtc ttt ttc cga gtg gca   288
     50 Ala Ala Val Asp Thr Asp Ser Pro Arg Glu Val Phe Phe Arg Val Ala
     51                  85                  90                  95
     53 gct gac atg ttt tct gac ggc aac ttc aac tgg ggc cgg gtt gtc gcc   336
     54 Ala Asp Met Phe Ser Asp Gly Asn Phe Asn Trp Gly Arg Val Val Ala
     55             100                 105                 110
     57 ctt ttc tac ttt gcc agc aaa ctg gtg ctc aag gcc ctg tgc acc aag   384
     58 Leu Phe Tyr Phe Ala Ser Lys Leu Val Leu Lys Ala Leu Cys Thr Lys
     59         115                 120                 125
     61 gtg ccg gaa ctg atc aga acc atc atg ggc tgg aca ttg gac ttc ctc   432
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(Cpg-6)

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PATENT APPLICATION: **US/10/550,280** TIME: 11:41:37

Input Set : **A:\Sequence Listing 2005_1500A.txt**
Output Set: **N:\CRF4\10062005\J550280.raw**

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62 Val Pro Glu Leu Ile Arg Thr Ile Met Gly Trp Thr Leu Asp Phe Leu
63     130                 135                 140
65 cgg gag cgg ctg ttg ggc tgg atc caa gac cag ggt ggt tgg gac ggc   480
66 Arg Glu Arg Leu Leu Gly Trp Ile Gln Asp Gln Gly Gly Trp Asp Gly
67 145                 150                 155                 160
69 ctc ctc tcc tac ttt ggg acg ccc acg tgg cag acc gtg acc atc ttt   528
70 Leu Leu Ser Tyr Phe Gly Thr Pro Thr Trp Gln Thr Val Thr Ile Phe
71                 165                 170                 175
73 gtg gcg gga gtg ctc acc gcc tcg ctc acc atc tgg aag aag atg ggc   576
74 Val Ala Gly Val Leu Thr Ala Ser Leu Thr Ile Trp Lys Lys Met Gly
75             180                 185                 190
77 tga                                                              579
80 <210> SEQ ID NO: 2
81 <211> LENGTH: 192
82 <212> TYPE: PRT
83 <213> ORGANISM: Homo sapiens
85 <400> SEQUENCE: 2
86 Met Asp Gly Ser Gly Glu Gln Pro Arg Gly Gly Gly Pro Thr Ser Ser
87   1               5                  10                  15
88 Glu Gln Ile Met Lys Thr Gly Ala Leu Leu Leu Gln Gly Phe Ile Gln
89              20                  25                  30
90 Asp Arg Ala Gly Arg Met Gly Gly Glu Ala Pro Glu Leu Ala Leu Asp
91          35                  40                  45
92 Pro Val Pro Gln Asp Ala Ser Thr Lys Lys Leu Ser Glu Cys Leu Lys
93      50                  55                  60
94 Arg Ile Gly Asp Glu Leu Asp Ser Asn Met Glu Leu Gln Arg Met Ile
95  65                  70                  75                  80
96 Ala Ala Val Asp Thr Asp Ser Pro Arg Glu Val Phe Phe Arg Val Ala
97                  85                  90                  95
98 Ala Asp Met Phe Ser Asp Gly Asn Phe Asn Trp Gly Arg Val Val Ala
99             100                 105                 110
100 Leu Phe Tyr Phe Ala Ser Lys Leu Val Leu Lys Ala Leu Cys Thr Lys
101         115                 120                 125
102 Val Pro Glu Leu Ile Arg Thr Ile Met Gly Trp Thr Leu Asp Phe Leu
103     130                 135                 140
104 Arg Glu Arg Leu Leu Gly Trp Ile Gln Asp Gln Gly Gly Trp Asp Gly
105 145                 150                 155                 160
106 Leu Leu Ser Tyr Phe Gly Thr Pro Thr Trp Gln Thr Val Thr Ile Phe
107                 165                 170                 175
108 Val Ala Gly Val Leu Thr Ala Ser Leu Thr Ile Trp Lys Lys Met Gly
109             180                 185                 190
113 <210> SEQ ID NO: 3
114 <211> LENGTH: 999
115 <212> TYPE: DNA
116 <213> ORGANISM: Homo sapiens
118 <220> FEATURE:
119 <221> NAME/KEY: CDS
120 <222> LOCATION: (1)..(999)
122 <400> SEQUENCE: 3
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RAW SEQUENCE LISTING DATE: 10/06/2005
PATENT APPLICATION: **US/10/550,280** TIME: 11:41:37

Input Set : **A:\Sequence Listing 2005_1500A.txt**
Output Set: **N:\CRF4\10062005\J550280.raw**

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123 atg gcc tgc gat tgc cgt ggt gat tgt ttt tgt ggt ggt atg agc aag   48
124 Met Ala Cys Asp Cys Arg Gly Asp Cys Phe Cys Gly Gly Met Ser Lys
125   1               5                  10                  15
127 ggc gag gaa ctg ttc act ggc gtg gtc cca att ctc gtg gaa ctg gat   96
128 Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu Asp
129             20                  25                  30
131 ggc gat gtg aat ggg cac aaa ttt tct gtc agc gga gag ggt gaa ggt   144
132 Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu Gly Glu Gly
133         35                  40                  45
135 gat gcc aca tac gga aag ctc acc ctg aaa ttc atc tgc acc act gga   192
136 Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys Thr Thr Gly
137     50                  55                  60
139 aag ctc cct gtg cca tgg cca aca ctg gtc act acc ttc acc tat ggc   240
140 Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe Thr Tyr Gly
141 65                  70                  75                  80
143 gtg cag tgc ttt tcc aga tac cca gac cat atg aag cag cat gac ttt   288
144 Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln His Asp Phe
145                 85                  90                  95
147 ttc aag agc gcc atg ccc gag ggc tat gtg cag gag aga acc atc ttt   336
148 Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe
149             100                 105                 110
151 ttc aaa gat gac ggg aac tac aag acc cgc gct gaa gtc aag ttc gaa   384
152 Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu
153         115                 120                 125
155 ggt gac acc ctg gtg aat aga atc gag ctg aag ggc att gac ttt aag   432
156 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys
157     130                 135                 140
159 gag gat gga aac att ctc ggc cac aag ctg gaa tac aac tat aac tcc   480
160 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser
161 145                 150                 155                 160
163 cac aat gtg tac atc atg gcc gac aag caa aag aat ggc atc aag gtc   528
164 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val
165                 165                 170                 175
167 aac ttc aag atc aga cac aac att gag gat gga tcc gtg cag ctg gcc   576
168 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val Gln Leu Ala
169             180                 185                 190
171 gac cat tat caa cag aac act cca atc ggc gac ggc cct gtg ctc ctc   624
172 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu
173         195                 200                 205
175 cca gac aac cat tac ctg tcc acc cag tct gcc ctg tct aaa gat ccc   672
176 Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser Lys Asp Pro
177     210                 215                 220
179 aac gaa aag aga gac cac atg gtc ctg ctg gag ttt gtg acc gct gct   720
180 Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val Thr Ala Ala
181 225                 230                 235                 240
183 ggg atc aca cat ggc atg gac gag ctg tac aag gcc ctt ttc tac ttt   768
184 Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys Ala Leu Phe Tyr Phe
185                 245                 250                 255
187 gcc agc aaa ctg gtg ctc aag gcc ctg tgc acc aag gtg ccg gaa ctg   816
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RAW SEQUENCE LISTING DATE: 10/06/2005
PATENT APPLICATION: **US/10/550,280** TIME: 11:41:37

Input Set : **A:\Sequence Listing 2005_1500A.txt**
Output Set: **N:\CRF4\10062005\J550280.raw**

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188 Ala Ser Lys Leu Val Leu Lys Ala Leu Cys Thr Lys Val Pro Glu Leu
189             260                 265                 270
190 atc aga acc atc atg ggc tgg aca ttg gac ttc ctc cgg gag cgg ctg   864
191 Ile Arg Thr Ile Met Gly Trp Thr Leu Asp Phe Leu Arg Glu Arg Leu
192         275                 280                 285
194 ttg ggc tgg atc caa gac cag ggt ggt tgg gac ggc ctc ctc tcc tac   912
195 Leu Gly Trp Ile Gln Asp Gln Gly Gly Trp Asp Gly Leu Leu Ser Tyr
196     290                 295                 300
198 ttt ggg acg ccc acg tgg cag acc gtg acc atc ttt gtg gcg gga gtg   960
199 Phe Gly Thr Pro Thr Trp Gln Thr Val Thr Ile Phe Val Ala Gly Val
200 305                 310                 315                 320
202 ctc acc gcc tca ctc acc atc tgg aag aag atg ggc tga               999
203 Leu Thr Ala Ser Leu Thr Ile Trp Lys Lys Met Gly
204                 325                 330
207 <210> SEQ ID NO: 4
208 <211> LENGTH: 332
209 <212> TYPE: PRT
210 <213> ORGANISM: Homo sapiens
212 <400> SEQUENCE: 4
213 Met Ala Cys Asp Cys Arg Gly Asp Cys Phe Cys Gly Gly Met Ser Lys
214   1               5                  10                  15
215 Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu Asp
216              20                  25                  30
217 Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu Gly Glu Gly
218          35                  40                  45
219 Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys Thr Thr Gly
220      50                  55                  60
221 Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe Thr Tyr Gly
222  65                  70                  75                  80
223 Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Gln His Asp Phe
224                  85                  90                  95
225 Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe
226             100                 105                 110
227 Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu
228         115                 120                 125
229 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys
230     130                 135                 140
231 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser
232 145                 150                 155                 160
233 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val
234                 165                 170                 175
235 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val Gln Leu Ala
236             180                 185                 190
237 Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu
238         195                 200                 205
239 Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser Lys Asp Pro
240     210                 215                 220
241 Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val Thr Ala Ala
242 225                 230                 235                 240
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RAW SEQUENCE LISTING DATE: 10/06/2005
PATENT APPLICATION: US/10/550,280 TIME: 11:41:37

Input Set : A:\Sequence Listing 2005_1500A.txt
Output Set: N:\CRF4\10062005\J550280.raw

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243 Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys Ala Leu Phe Tyr Phe
244                 245                 250                 255
245 Ala Ser Lys Leu Val Leu Lys Ala Leu Cys Thr Lys Val Pro Glu Leu
246             260                 265                 270
247 Ile Arg Thr Ile Met Gly Trp Thr Leu Asp Phe Leu Arg Glu Arg Leu
248         275                 280                 285
249 Leu Gly Trp Ile Gln Asp Gln Gly Gly Trp Asp Gly Leu Leu Ser Tyr
250     290                 295                 300
251 Phe Gly Thr Pro Thr Trp Gln Thr Val Thr Ile Phe Val Ala Gly Val
252 305                 310                 315                 320
253 Leu Thr Ala Ser Leu Thr Ile Trp Lys Lys Met Gly
254                 325                 330
258 <210> SEQ ID NO: 5
259 <211> LENGTH: 987
260 <212> TYPE: DNA
261 <213> ORGANISM: Homo sapiens
263 <220> FEATURE:
264 <221> NAME/KEY: CDS
265 <222> LOCATION: (1)..(987)
267 <400> SEQUENCE: 5
268 atg gcc tgc aac ggt cgt tgc ggt ggt atg agc aag ggc gag gaa ctg   48
269 Met Ala Cys Asn Gly Arg Cys Gly Gly Met Ser Lys Gly Glu Glu Leu
270   1               5                  10                  15
272 ttc act ggc gtg gtc cca att ctc gtg gaa ctg gat ggc gat gtg aat   96
273 Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu Asp Gly Asp Val Asn
274             20                  25                  30
275 ggg cac aaa ttt tct gtc agc gga gag ggt gaa ggt gat gcc aca tac   144
276 Gly His Lys Phe Ser Val Ser Gly Glu Gly Glu Gly Asp Ala Thr Tyr
277         35                  40                  45
279 gga aag ctc acc ctg aaa ttc atc tgc acc act gga aag ctc cct gtg   192
280 Gly Lys Leu Thr Leu Lys Phe Ile Cys Thr Thr Gly Lys Leu Pro Val
281     50                  55                  60
283 cca tgg cca aca ctg gtc act acc ttc acc tat ggc gtg cag tgc ttt   240
284 Pro Trp Pro Thr Leu Val Thr Thr Phe Thr Tyr Gly Val Gln Cys Phe
285 65                  70                  75                  80
287 tcc aga tac cca gac cat atg aag cag cat gac ttt ttc aag agc gcc   288
288 Ser Arg Tyr Pro Asp His Met Lys Gln His Asp Phe Phe Lys Ser Ala
289                 85                  90                  95
291 atg ccc gag ggc tat gtg cag gag aga acc atc ttt ttc aaa gat gac   336
292 Met Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe Phe Lys Asp Asp
293             100                 105                 110
295 ggg aac tac aag acc cgc gct gaa gtc aag ttc gaa ggt gac acc ctg   384
296 Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu Gly Asp Thr Leu
297         115                 120                 125
299 gtg aat aga atc gag ctg aag ggc att gac ttt aag gag gat gga aac   432
300 Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys Glu Asp Gly Asn
301     130                 135                 140
303 att ctc ggc cac aag ctg gaa tac aac tat aac tcc cac aat gtg tac   480
304 Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser His Asn Val Tyr
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 10/06/2005
PATENT APPLICATION: **US/10/550,280** TIME: 11:41:38

Input Set : **A:\Sequence Listing 2005_1500A.txt**
Output Set: **N:\CRF4\10062005\J550280.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:19; N Pos. 1,2
Seq#:22; N Pos. 1,2
Seq#:24; N Pos. 1,2,3,4
Seq#:27; N Pos. 1,2,3,4

VERIFICATION SUMMARY DATE: 10/06/2005
PATENT APPLICATION: **US/10/550,280** TIME: 11:41:38

Input Set : **A:\Sequence Listing 2005_1500A.txt**
Output Set: **N:\CRF4\10062005\J550280.raw**

L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:420 M:283 W: Missing Blank Line separator, <220> field identifier
L:448 M:283 W: Missing Blank Line separator, <400> field identifier
L:476 M:283 W: Missing Blank Line separator, <400> field identifier
L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:609 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:636 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:667 M:283 W: Missing Blank Line separator, <220> field identifier
L:673 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0